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TITLE: Eicosanoid Regulation of Prostate Cancer Progression: Disruption of Hemidesmosomes and Collaboration in Tumor Invasive Growth

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INTRODUCTION

During the progression of human PCa hemidesmosomes, adhesion structures that anchor epithelial cells to basement membrane and function as a tumor suppressor, are lost (1, 2). We found that 12lipoxygenase directly interacts with $\beta4$ integrin, an integral part of hemidesmosomes (3, 4). We hypothesized that an increase in 12activity can cause the disassembly of hemidesmosomes, mobilization of $\alpha 6\beta 4$ integrin from hemidesmosomes to other parts of the cell membrane, and stimulate tumor invasive growth. evaluate proposed to conduct а correlation study to 12-LOX expression and dispersion of relationship between integrin in clinical tumor specimens. Our proposal further aims to study whether 12(S)-HETE, the enzymatic product of 12-LOX, can disrupt hemidesmosomes and whether 12-LOX inhibitors promote the formation of hemidesmosomes. Then we will study the underlying signaling pathway, especially $PKC\alpha$, initiated by 12(S)-HETE, in the disassembly of hemidesmosomes. Next we will overexpress $\beta 4$ integrin and study the role of the interaction between 12-LOX and integrin in the adhesion, proliferation, migration, survival, in response to HGF/SF. Finally we will xenograft these transfected cells into mice, to evaluate whether any phenotypic changes of tumor cells in vitro can be recapitulated in vivo. The work will significantly advance our understanding about complex process of prostate cancer progression as well as the possible role played by dietary fat in the progression of prostate cancer.

BODY OF PROGRESS REPORT

Task 1.

We have attempted several procedures for immunostaining for 12-LOX. In the previous report, the procedure only worked in frozen human prostate tumor tissue. We now worked out the conditions for immunohistochemical analysis of 12-LOX at the protein level in paraffin-embedded human prostate tumor tissues. As shown in the figure 1, 12-LOX immunoreactivity correlated with tumor grade. Neoplastic glands are weakly, moderately or strongly positive for 12-LOX in hyperplatic glands (Figure 1A), atropic glands (Figure 1B), and in tumor Glenson score 4-6 (Figure 1 C,D).

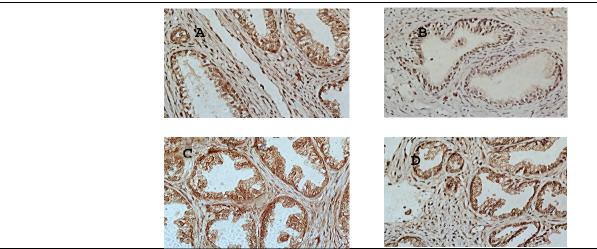


Figure 1. Immunohistochemical staining for 12-LOX in parafirm-embedded prostate tumor tissues. Sections of specimens were probed with a12-LOX polyclonal antibody (Oxford Biomedical Research Inc, Oxford, MI). Positive immunoreactivity is indicated by staining with brownish color. A, hyperplasia gland; B, atropic; C/D, tumor with Glenson score 4-6.

We have also attempted several protocols of immunostaining for $\beta 4$ integrin in paraffin-embedded material. As shown in figure 2, positive staining was found in tumors and correlated with tumor grade.

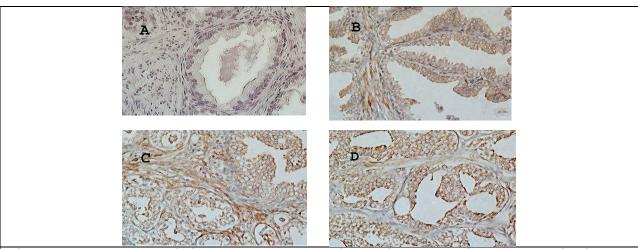


Figure 2. Immunostaining for beta 4 integrin. Brownish color indicates positive staining. Note the pattern of beta 4 integrin positivity. A, normal epithelium; B, tumor with Glenson score 4; C, tumor with Glenson score 6; D, tumor with Glenson score 8.

We have procured 100 cases of prostate tumor specimens. We are working on the protocol for double staining, the study will be completed in the next year.

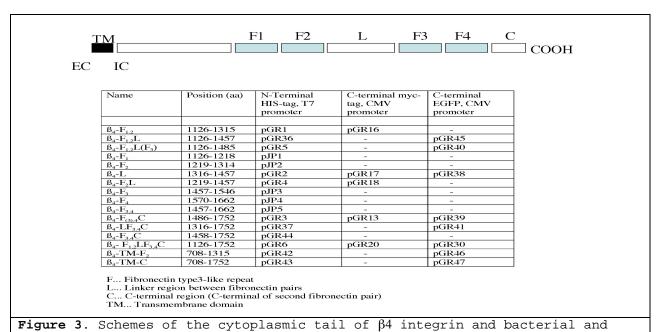
Task 2. Study the effects of 12-LOX inhibitors and 12(S)-HETE on hemidesmosome in prostate epithelium. Months 1-18: This specific aim was partially achieved during the previous reporting period and it is in the final stages.

Task 3. Study the signal transduction pathways that underlie the disassembly of hemidesmosomes by 12(S)-HETE or an increase in 12-LOX activity, Months 12 -24:

The studies proposed have been initiated and are ongoing.

Task 4. Overexpress $\beta 4$ integrin in PC-3 cells, in the presence or absence of 12-LOX expression, and evaluate the capacity of transfected cells to form hemidesmosomes and whether an increase in surface expression of $\alpha 6\beta 4$ alters cell proliferation, adhesion, migration, and survival, in response to HGF/SF, Months 18 - 30:

Generation of PC-3 cells expressing various $\beta4$ mutants: We constructed a panel of bacterial expression plasmids and mammalian expression constructs that express various $\beta4$ mutants of the cytoplasmic tail as shown in Figure 3. Using pGR30~47, we have generated a panel of stable transfectants that express various $\beta4$ mutants as GFP fusion proteins through fluorescence activated

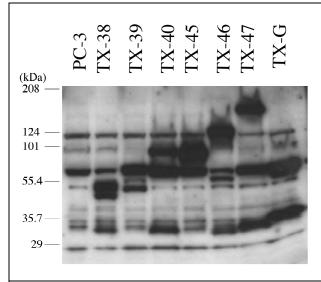


mammalian expression constructs. TM, transmembrane; EC, extracellular; IC, intracellular

cell sorting (FACS) and G418 selection for the transfectants.

The expression of $\beta4$ mutants as GFP-fusion proteins is shown in figure 4. Co-immunoprecipitation was performed to determine the interaction of 12-LOX with various $\beta4$ mutants. As shown in the

figure 5, 12-LOX is able to bind to the fusion proteins expressed by pGR40, pGR45, pGR46, and pGR47, but not to the fusion protein with only the linker region (pGR38). 12-LOX weakly binds to the fusion protein encoded by pGR39. Collectively, the data suggest a strong binding site(s) for 12-LOX located between 1126-1315 of $\beta 4$ cytoplasmic tail.



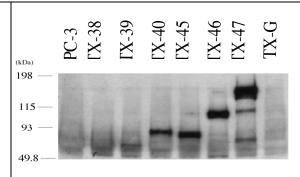
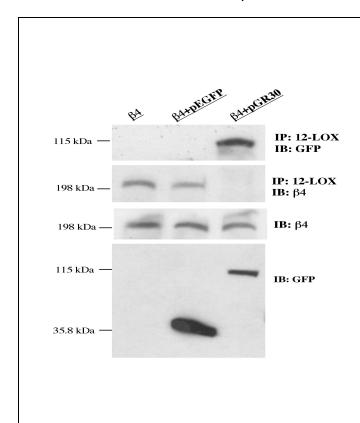


Figure 4. Western blot analysis for Figure 5. Co-immunoprecipitation of the expression of β 4 mutants in PC-3 | 12-LOX with the transfectants with an antibody against GFP. Note the expression of GFP fusion proteins of various MWs. calculated MW for fusion proteins for pGR47 (Tx47) is about 146. For pGR46 (Tx46), 97.4; pGR45 (Tx45), 66.8; pGR40 (Tx40), 70; pGR39 (Tx39), 59.6; and pGR38 (Tx38), 45.7. The MW for pEGFP (TxG) is approximately 30.

fusion proteins expressed by pGR40, pGR45, pGR46, and pGR47, but not to pGR38 and pEGFP (TxG). It is deduced that a binding site or sites are located between 1126 - 1315 region of cytoplasmic tail.

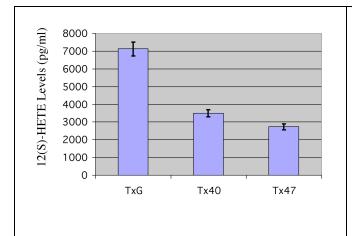
Cytosolic β 4 cytoplasmic tail binds to 12-LOX and blocks interaction between 12-LOX and full length $\beta 4$ integrin: As shown above, we have constructed a panel of expression constructs encoding various mutants for the $\beta4$ cytoplasmic tail. The construct pGR30 encodes the cytosolic $\beta4$ cytoplasmic tail, with TM domain deleted, as a GFP-fusion protein (Figure 3). When ectopically expressed in 12-LOX transfected PC-3 cells, the cytosolic $\beta4$ cytoplasmic tail interacts with 12-LOX (Figure 6, the IP: 12-LOX / IB: GFP panel). Interestingly the ectopically expressed cytosolic $\beta4$ tail blocks the interaction of 12-LOX with full-length $\beta4$ integrin (Figure 6, the IP: $12-LOX / IB: \beta 4$ panel). The blockade of the interaction between 12-LOX and full-length β4 integrin is not due to the lack of β 4 expression (Figure 6, the IB: β 4 panel), but due to the presence of the cytosolic $\beta 4$ cytoplasmic tail (Figure 6, the IB: GFP panel). The results suggest that the cytosolic β 4 mutant, encoded by pGR30, is able to block the interaction between 12-LOX and full-length $\beta4$ integrin in a dominant negative manner.



6: Figure Cytosolic β4 encoded by pGR30, binds to 12-LOX blocks interaction the between 12-LOX and full length $\beta4$ integrin. 12-LOX transfected PC-3 cells (PC3/12LOX) transiently transfected with an expression construct for fulllength $\beta 4$, in the presence or absence of pGR30 or its vector, Cell lysates pEGFP. immunoprecipitated with а polyclonal antibody against 12-LOX and immunoblotted for GFP Clontech) or full-(JL8 mAb, β4 integrin (AB1922, Chemicon International. Target band, around 205 kDa). Total lysates were also analyzed for the expression of full-length $\beta 4$ integrin and GFP or GFP-fusion protein by immunoblot. Note the co-immunoprecipitation of 12-LOX and the cytosolic $\beta4$ cytoplasmic tail and the blockade of interaction between 12-LOX full-length $\beta4$ integrin by cotransfection with pGR30.

Task 5. Evaluate the growth rates of s.c. tumors derived from $\alpha6\beta4$ expressing PC-3 cells, in the presence or absence of stable 12-LOX expression, and compare with that of control PC-3 cells, Months 24-36:

We have conducted a preliminary study to determine whether ectopic expression of $\beta4$ mutants, especially those that can bind 12-LOX, can effect 12-LOX activity. As shown in Figure 7, there in 12(S)-HETE biosynthesis in PC-3 cells reduction expressing two 12-LOX binding $\beta4$ mutants (pGR40 and pGR47). We are in the process of determining whether 12-LOX cellular localization and interaction with the full-length $\beta4$ integrin are altered as result of the presence of $\beta4$ mutants encoded by pGR40 and pGR47. When injected into athymic nu/nu mice, it was found, as shown in Figure 8, that there were a reduction in tumor growth rate in PC-3 cells expressing 12-LOX binding $\beta4$ mutants (Tx40 and Tx47, see figure 5), when compared to those from vector control (TxG) or a β 4 fragment that does not interact with 12-LOX (Tx38, see Figure 5).



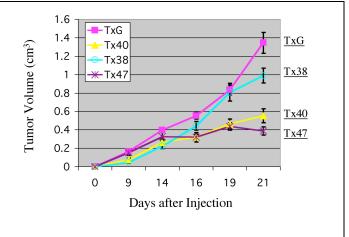


Figure 7. Decrease in 12(S)-HETE biosynthesis in PC-3 cells that express 12-LOX binding $\beta4$ mutants. 12(S)-HETE was isolated from cell lystaes and measured by EIA. N = 6 for each type of cells. Bar, SEM.

8. Alteration Figigure in growth as a result of ectopic expression of 12-LOX binding $\beta4$ mutants (Tx40 and Tx47), not non-12LOX-bind mutant (Tx38). TxG is the vector control. Each datum point, the average tumor volume from 4 mice. Bar, SEM. Note the correlation of reduction in tumor growth reduced 12(S)-HETE biosynthesis shown in figure 7.

Key Research Accomplishments:

- Optimized the conditions for immunohistochemical analysis of 12-LOX and $\beta4$ at the protein level in paraffin-embedded human prostate tumor tissues. This standardized method will enable us to continue with the correlation between 12-LOX expression and distribution of $\beta4$ integrin with Gleason score.
- Constructed several $\beta4$ cytoplasmic tail mutants and identified the sequence of amino acids on $\beta4$ integrin that interacts with 12-LOX.
- Preliminary experiments conducted to show the feasibility of disruption of the interaction of 12-LOX with $\beta4$ integrin in PC-3 cells with ectopic expression of the cytoplasmic tail sequence of $\beta4$ integrin. This resulted in the reduction of the biosynthesis of 12-HETE as well as tumor growth from PC-3 cells in experimental animals.

Reportable Outcomes:

NONE

Conclusions:

Interaction of 12-LOX with the cytoplasmic tail of $\beta 4$ integrin results in the disruption of hemidesmosomes. Our data presented here demonstrates the potential to exploit the interaction between

12-LOX and $\beta 4$ integrin using ectopically expressed cytoplasmic tail sequence of the integrin to modulate tumor growth.

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Appendices:

NONE